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Regulation of the DNA damage response on male meiotic sex chromosomes

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Abstract

During meiotic prophase in males, the sex chromosomes partially synapse to form the XY body. This is a unique structure that recruits proteins involved in the DNA damage response, which is believed to be important for silencing of the sex chromosomes. It remains elusive how the DNA damage response in the XY body is regulated. H2AX-MDC1-RNF8 signaling, which is well characterized in somatic cells, is dispensable for the recruitment of proteins to the unsynapsed axes in the XY body. However, the DNA damage response that spreads over the sex chromosomes is largely similar to that in somatic cells. Here we show that accumulation of some components of the DNA damage response pathway on the XY body occurs upstream of H2AX-MDC1-RNF8 signalling, and downstream from this cascade of events for others. This analysis shows that there are important differences between the regulation of the DNA damage response at the XY body and at DNA damage sites in somatic cells.

Introduction

Meiotic prophase is one of the most important stages before meiotic divisions in both male and female reproductive germ cells. During this period, homologous chromosomes synapse and undergo meiotic recombination. The exchange of genetic material between homologous chromosomes is critical for generating genetic diversity in the offspring. Meiotic recombination starts with global SPO11-dependent DNA breaks throughout the chromosomes, which trigger the DNA damage response (DDR)¹. Meiotic recombination occurs during chromosome synapsis, when DNA repair takes place through homologous recombination. One unique challenge during male meiosis is that male spermatocytes possess X and Y sex chromosomes, which are largely heterologous and only partially synapse through their pseudo autosomal regions (PAR). When autosomes are fully synapsed

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Author Contributions

L.L. and X.Y. conceived and designed the project. L.L., Y.X., H.K., and G.K. performed the experiments. L.L. and X.Y. prepared the manuscript. All authors discussed and commented on the manuscript.

Competing financial interests

The authors declare no competing financial interests.

and meiotic recombination is finished at pachynema of meiotic prophase, DDR proteins are lost from the autosomes and autosomes become transcriptionally active. However, the partially synapsed sex chromosomes at pachynema locate to a region close to the edge of the nucleus known as the XY body, and remain transcriptionally silenced. This phenomenon is known as meiotic sex chromosome inactivation (MSCI)². It is known that the XY body retains a lot of DDR proteins, which are believed to be important for the initiation of MSCI. Consistent with this idea, spermatocytes deficient for H2AX or MDC1, two proteins that act early in the DDR pathway in somatic cells, abolished MSCI^{3,4}.

Although many DDR proteins accumulate in the XY body, their localizations are different⁴. Some proteins, such as H2AX and MDC1, spread over the sex chromosomes. Some proteins, like ATR and TopBP1, are more concentrated at the unsynapsed axes, and are weakly stained on the sex chromosomes. Some proteins, including BRCA1, are exclusively located at the unsynapsed axes. In addition, single-stranded DNA binding protein RPA and DNA repair protein RAD51 form foci specifically at the unsynapsed axes. This suggests that the unsynapsed axes are different from other chromosomal regions of the XY body, and that the regulation of DDR in spermatocytes could be distinct from that in somatic cells. Recent studies have suggested that the DDR begins by recruiting some proteins to the unsynapsed axes in a MDC1-independent manner. This is then followed by MDC1-dependent amplification of the DDR to other regions of the sex chromosomes⁴.

RNF8 is a protein immediately downstream of MDC1 in the DDR in somatic cells^{5,6}. Different from *Mdc1* knockout spermatocytes that arrest at mid-pachynema without proper formation of the XY body, meiosis in *Rnf8* knockout mice progresses normally and the XY body forms readily in spermatocytes at pachynema⁷. Therefore, *Rnf8* knockout mice are better tools for examining the DDR signaling at the meiotic sex chromosomes in the XY body. In this study, we analyze the DDR signaling of the XY body in *Rnf8* knockout spermatocytes and explore potential mechanisms that recruit the DDR proteins to the unsynapsed axes. We find that there are important differences between the regulation of the DNA damage response at the XY body and at DNA damage sites in somatic cells.

Results

RNF8 controls DDR protein localization on sex chromosomes

Consistent with our prior observations, RNF8 deficiency has no obvious impact on meiotic prophase; and spermatocytes still progress through pachynema with normal XY body formation. RNF8 is known to be essential for protein ubiquitination in the XY body (Figure 1a)⁷. To distinguish which ubiquitin species is enriched in the XY body, we tested antibodies that specifically recognize K48 or K63-linked ubiquitin chains⁸. In agreement with the well characterized observation that RNF8 promotes the synthesis of K63-linked ubiquitin chains at DNA damage sites in somatic cells⁹, K63-linked ubiquitination was enriched in the XY body and was absent in *Rnf8* knockout spermatocytes (Figure 1b). Recent studies have shown that RNF8 also facilitates the transient assembly of K48-linked ubiquitin chains at DNA damage sites in somatic cells^{10,11}. Interestingly, K48-linked ubiquitination was also enriched in the XY body of wild type, but not *Rnf8* knockout spermatocytes (Figure 1c). These results suggest that RNF8 regulates both K48 and K63-

linked ubiquitination in the XY body. Besides ubiquitination, sumoylation is another ubiquitination-like protein modification at DNA damage sites^{12,13}. It has been shown that RNF8 is important for protein modification by SUMO1 or SUMO2/3 at DNA damage sites¹³, and both of them have been shown to enrich in the XY body and spread over the sex chromosome¹⁴. Interestingly, we found that both SUMO1 and SUMO2/3 still localized to the XY body in *Rnf8* knockout spermatocytes (Figure 1d–e). Therefore, RNF8 is dispensable for protein sumoylation in the XY body, which is different from the DDR in somatic cells.

Since RNF8 is important for propagating DNA damage signals and controls the localization of a number of proteins at DNA damage sites in somatic cells, we examined if RNF8 deficiency has any impact on the localization of a number of DDR proteins in spermatocytes. Proteins in somatic DDR pathways upstream of RNF8 (H2AX and MDC1) were still enriched in the XY body and spread over the sex chromosomes in *Rnf8* knockout spermatocytes (Figure 2a–b). It is well characterized that 53BP1 is downstream of RNF8 in the DDR of somatic cells^{5,6}. 53BP1 was localized at the XY body and spread over the sex chromosomes in wild type spermatocytes, but was absent in *Rnf8* knockout spermatocytes (Figure 2c). Although how RNF8 regulates 53BP1 in somatic DDR is not clear, it seems that it regulates 53BP1 using the same mechanism in spermatocytes. It is also known that the RNF8-dependent ubiquitin chains directly recruit RAD18 to DNA damage sites in somatic cells¹⁵, and RAD18 spreads over the sex chromosomes¹⁶. Interestingly, the localization of RAD18 to the XY body was completely abolished in *Rnf8* knockout spermatocytes, which is consistent with the absence of ubiquitin chains (Figure 2d). Since the absence of both 53BP1 and RAD18 in the XY body does not affect the meiosis of *Rnf8* knockout spermatocytes, these two proteins are likely dispensable for meiosis, which is consistent with previous studies of mice lacking either of them^{17–19}. Besides proteins that spread over the sex chromosomes, some proteins in the XY body, like ATR and TopBP1, spread over the sex chromosomes with enrichment in the unsynapsed axes²⁰. In agreement with the observation that RNF8 deficiency does not affect ATR or TopBP1 foci in somatic cells²¹, there was little difference in the localizations of ATR or TopBP1 between wild type and *Rnf8* knockout spermatocytes (Figure 2e–f). Although it has been suggested that MDC1 is required for the spread of ATR and TopBP1 over the sex chromosomes⁴, RNF8 seems to be dispensable for this process.

Localization of the BRCA1-A complex does not rely on RNF8

Similar to RAD18, RAP80 also localizes to DNA damage sites through recognition of RNF8-dependent ubiquitin chains in somatic cells^{22–24}. But unlike RAD18 that spreads over the sex chromosome, RAP80 is localized specifically to the unsynapsed axes in wild type spermatocytes (Figure 3a). Interestingly, the localization of RAP80 was still preserved in the *Rnf8* knockout spermatocytes (Figure 3a). These results suggest that the regulation of RAD18 and RAP80 is different in spermatocytes. RAP80 belongs to a big complex containing BRCA1 in somatic cells known as the BRCA1-A complex^{25–27}, and RAP80 is required for targeting this complex to the DNA damage sites in somatic cells. To examine if RAP80 belongs to the same BRCA1-A complex in spermatocytes, we determined the localizations of other members of this complex, including BRCA1, CCDC98, and

MERIT40, in spermatocytes. Consistent with our previous observation, BRCA1 localized specifically to the unsynapsed axes, and the localization was not affected by RNF8 deficiency (Figure 3b). Interestingly, both CCDC98 and MERIT40 also localized to the unsynapsed axes in both wild type and *Rnf8* knockout spermatocytes (Figure 3c–d). Although we did not check the localizations of BRCC36 and BRCC45, due to unavailability of antibodies, our observations suggest that the same BRCA1-A complex exists in spermatocytes and its members localize at the unsynapsed axes in spermatocytes. Unlike the DDR in somatic cells, the localization of BRCA1-A complex members is independent of RNF8. This suggests that the BRCA1-A complex does not recognize RNF8-dependent ubiquitin chains in spermatocytes by RAP80, as it does during DDR in somatic cells. The BRCA1-A complex is known to promote homologous recombination repair by recruiting RAD51 to DNA damage sites in somatic cells²⁸. Consistent with the localization of the BRCA1-A complex, RAD51 was exclusively located at the unsynapsed axes (Figure 3e). Interestingly, although RNF8 regulates the localization of RAD51 to DNA damage sites in somatic cells²⁹, it had no impact on the localization of RAD51 at the unsynapsed axes in spermatocytes (Figure 3e). These observations suggest that RNF8 is dispensable for homologous recombination repair of the sex chromosomes, which is likely mediated by the BRCA1-A complex and RAD51.

Since the BRCA1-A complex is also regulated by H2AX and MDC1, two proteins upstream of RNF8 in the DDR in somatic cells, we wondered if the BRCA1-A complex is still regulated by H2AX and MDC1 in spermatocytes. Interestingly, BRCA1 still localized at the unsynapsed axes in *H2ax* and *Mdc1* knockout spermatocytes (Figure 4a). Moreover, the localizations of RAP80, CCDC98, and MERIT40 at unsynapsed axes were also preserved in *H2ax* and *Mdc1* knockout spermatocytes (Figure 4b–d). These observations suggest that the accumulation of the BRCA1-A complex at the unsynapsed axes occurs through other mechanisms.

BRCA1's localization relies on BRCT domain

To gain more insight into the mechanism by which BRCA1 localizes to the unsynapsed axes in spermatocytes, we examined if it localizes through its BRCT domains, as it does at DNA damage sites in somatic cells³⁰. The BRCA1 BRCT domain is a phospho-binding module³⁰. To study if the BRCA1 BRCT domain could still function as a phospho-binding module in spermatocytes, we developed a novel *in vitro* binding method to examine the protein localization in spermatocytes. Recombinant GST-tagged BRCA1 BRCT domain was incubated with the spermatocyte spread, and the localization of the BRCA1 BRCT domain in spermatocytes was then visualized by immunostaining using an anti-GST antibody. While GST alone, a negative control, could not be observed in spermatocytes (Figure 5a), the BRCA1 BRCT domain clearly localized at the unsynapsed axes (Figure 5b). In addition, the localization of the recombinant BRCA1 BRCT domain at unsynapsed axes could be observed in *H2ax* knockout, *Mdc1* knockout, and *Rnf8* knockout spermatocytes (Figure 5c). This fully recapitulated the pattern of endogenous BRCA1 in these cells. To examine if the localization of BRCA1 requires the presence of phosphorylated proteins at the unsynapsed axes, we pretreated the spermatocyte spread with lambda phosphatase before incubating it with the recombinant BRCA1 BRCT domain. Lambda phosphatase specifically abolished

the protein phosphorylation at the unsynapsed axes, including CCDC98 phosphorylation, after the pretreatment (Figure 5d–e). Although RAP80, which localizes to the unsynapsed axes in a phosphorylation-independent manner, remained at the unsynapsed axes (Figure 5f), the recombinant BRCA1 BRCT domain could no longer localize to the unsynapsed axes following phosphatase treatment (Figure 5g). Consistent with this observation, mutations of the key amino acids in the phospho-binding pocket of BRCA1 BRCT domain (S1655A or K1702A) completely abolished its localization to the unsynapsed axes (Figure 5h). Together, these results suggest that the recruitment of BRCA1 to the unsynapsed axes is indeed through binding of phosphorylated proteins using its BRCT domains.

It has been shown that phospho-dependent binding of CCDC 98 in the BRCA1-A complex is required for targeting BRCA1 to DNA damage sites in somatic cells^{22,31,32}. We wondered if the same mechanism is used for targeting BRCA1 to the unsynapsed axes. We examined mice deficient for RAP80, which is known to be required for targeting the BRCA1-A complex to DNA damage sites through recognition of protein ubiquitination in somatic cells³³. *Rap80* knockout mice are fertile and display no defect in meiotic progression. Analysis of *Rap80* knockout spermatocytes revealed that CCDC98 is absent from the unsynapsed axes of the XY body (Figure 6a), but interestingly BRCA1 is still preserved (Figure 6b). This suggests that the BRCA1-A complex is dispensable for targeting BRCA1 to the unsynapsed axes. It is possible that BRCA1 is targeted to the unsynapsed axes by other proteins, and that the BRCA1-A complex is tethered there through BRCA1. This is consistent with the observation that loss of ubiquitination in *Rnf8* knockout spermatocytes does not affect the localization of the BRCA1-A complex (Figure 3a–d). To search for other mechanisms for targeting BRCA1 to the unsynapsed axes in a phospho-dependent manner, we analyzed the localizations of two other known somatic BRCA1 phospho-dependent binding partners: BACH1 from the BRCA1-B complex³⁰, and CTIP from the BRCA1-C complex^{34,35}. Interestingly, although BACH1 was not detected in the XY body (Figure 6c), CTIP was specifically localized to the unsynapsed axes of the XY body (Figure 6d). The localization of CTIP was not altered in *H2ax*, *Mdc1*, *Rnf8*, or *Rap80* knockout mice (Figure 6e), which fully coincide with the localizations of BRCA1 in these mice. Since direct evaluation of the impact of the loss of CTIP on the localization of BRCA1 could not be performed due to the embryonic lethality of *Ctip* knockout mice³⁶, it is not clear if CTIP plays any roles in targeting BRCA1 to the unsynapsed axes of the XY body.

ATM and DNAPK localize to unsynapsed axes of sex chromosomes

Since protein phosphorylation is important for BRCA1 localization to the unsynapsed axes independent of H2AX-MDC1-RNF8 signaling, we wondered which protein kinases are responsible for the protein phosphorylation located at the unsynapsed axes. ATR is a protein kinase that is known to localize to the unsynapsed axes⁴. Besides ATR, ATM and DNAPK are two other important protein kinases for the DDR in somatic cells and are autophosphorylated when they are activated. We examined the localizations of the active forms of these two kinases and found that both of them localized at the unsynapsed axes (Figure 7a–b). Interestingly, the localizations of all these three protein kinases were still preserved in spermatocytes lacking H2AX, MDC1, or RNF8 (Figure 7a–b). Since the activations of ATM and DNAPK at DNA damage sites require proper H2AX-MDC1-RNF8

signaling in somatic cells^{37,38}, this observation suggests that the regulation of these kinases at the unsynapsed axes is different from that in somatic DDR. Future studies will reveal their precise roles at the unsynapsed axes.

Discussion

Many proteins in the DDR pathways accumulate in the XY body of spermatocytes at pachynema. This suggests that the DNA breaks are not yet repaired on the sex chromosomes. The delay of the repair might be caused by the fact that the sex chromosomes do not have homologous chromosomes, which are favorable templates for homologous recombination repair during meiotic prophase. The DNA breaks on the sex chromosomes are likely to be repaired by homologous recombination using their sister chromatids instead, which is not preferred and used as a back-up at the later stage of meiotic prophase. During this inefficient repair process in the XY body, transcription on the sex chromosomes is silenced, which is known as meiotic sex chromosome inactivation (MSCI)². Although the mechanism is not clear, it is likely that MSCI is similar to DNA damage-induced transcriptional silencing in somatic cells³⁹, which is a protective measure downstream of the DDR to prevent the generation of truncated RNA.

Although many DDR proteins accumulate in the XY body, it is puzzling why some proteins are restricted to the unsynapsed axes while others spread over the sex chromosomes. This unique feature might be a result of the prolonged presence of DNA damage in the XY body. It has been shown that the localization of proteins to the unsynapsed axes is independent of canonical H2AX-MDC1 signaling^{3,4}. In this study, we have identified several new proteins that are restricted to the unsynapsed axes, and their localization is not regulated by H2AX-MDC1-RNF8 pathways either. This is reminiscent of the observations in somatic cells that the localization of some proteins to the DNA damage sites at early time points is independent of H2AX, MDC1, or RNF8^{40,41}. It is therefore possible that similar mechanisms for protein recruitment are shared between these two events. Previous studies have shown that ATR is localized at the unsynapsed axes. Interestingly, we observed the localization of two other DDR-related protein kinases (ATM and DNAPK) at the unsynapsed axes. This suggests that protein phosphorylation is indeed important for the DDR signaling at unsynapsed axes. Recent studies suggest that these protein kinases at the unsynapsed axes are also regulated by proteins containing the HORMA domain, which preferentially associate with unsynapsed chromosomes and are likely to contribute to the initiation of the DDR on the unsynapsed axes of the XY body⁴²⁻⁴⁶. In agreement with this, BRCA1 is likely to be recruited through binding to phosphorylated proteins. Since BRCA1 is upstream of the localization of ATR to unsynapsed axes^{47,48}, it is likely that ATM and/or DNAPK-dependent protein phosphorylation recruits BRCA1 to the unsynapsed axes. The identity of the phosphorylated protein remains to be found.

Besides the proteins enriched at the unsynapsed axes, other DDR proteins spread over the sex chromosomes. This is similar to the observation in somatic cells that DDR proteins spread megabases away from the actual DNA breaks⁴⁹. The spread of DDR proteins over the sex chromosomes explains why the whole sex chromosomes, but not only the regions at the unsynapsed axes, are transcriptionally silenced. Based on the observation that DDR

proteins no longer spread over the sex chromosomes in *Mdc1* knockout spermatocytes, it was suggested that H2AX and MDC1 control the spread of DDR proteins over the whole sex chromosomes despite the fact that they are dispensable for the signaling on the unsynapsed axes⁴. However, since *Mdc1* knockout spermatocytes arrest at mid-pachynema without proper formation of the XY body, it is difficult to distinguish if the defective spreading of DDR proteins to the sex chromosomes is due to the absence of MDC1's function in the DDR signaling or the absence of XY body where the spreading takes place. In this respect, *Rnf8* knockout spermatocytes are more suitable for studying the DDR signaling over the sex chromosomes, since there is no defect in meiotic progression and the XY body readily forms at pachynema in *Rnf8* knockout spermatocytes. Using *Rnf8* knockout spermatocytes, we found that the DDR signaling of the sex chromosome is largely similar to the DDR signaling in somatic cells. The proteins upstream of RNF8 (H2AX and MDC1) are retained and those downstream of RNF8 (K63-Ub, K48-Ub, 53BP1, and RAD18) are absent in *Rnf8* knockout spermatocytes. Differences were also noticed between the signaling pathways in spermatocytes and somatic cells. For example, protein sumoylation is not dependent on RNF8 in spermatocytes, which is consistent with the observation that sumoylation precedes H2AX phosphorylation in the XY body⁵⁰. It has been shown that the modifications by SUMO1 and SUMO2/3 mainly occur on BRCA1 and 53BP1, respectively, during the DDR in somatic cells^{12,13}. Since BRCA1 does not localize over the whole sex chromosomes and the spread of 53BP1 over sex chromosomes is absent in *Rnf8* knockout spermatocytes, it is likely that the modifications by SUMO1 and SUMO2/3 on sex chromosomes predominantly occur on other proteins, which remain to be identified in spermatocytes.

In summary, we have studied the regulation of DDR signaling in the XY body of spermatocytes at pachynema. In our model, some proteins are recruited to the unsynapsed axes independent of canonical DDR pathways. The DDR then spreads over the whole sex chromosomes by amplification of γ H2AX-MDC1-RNF8 signaling, which recruits another set of proteins (Figure 8). We have also found that *Rnf8* knockout mice are invaluable tools for studying meiotic DDR. In addition, we have generated a novel *in vitro* binding method for analyzing DDR proteins, which allows easy visualization of DDR proteins at the XY body while circumventing the need for antibodies against these proteins.

Material and methods

Mice

Animal studies were ethically approved by the University Committee on Use and Care of Animals (UCUCA) of the University of Michigan. *Rnf8* and *Rap80* knockout mice were reported previously^{7,33}. *H2ax* and *Mdc1* knockout mice are from Andre Nussenzweig and Zhenkun Lou, respectively. All of the mice were housed in an AAALAC accredited facility in accordance with the National Research Council's guide for the care and use of laboratory animals.

Immunofluorescence staining

Immunofluorescence staining (IF) was performed at room temperature. The slides were first blocked with IF buffer (5% goat serum in PBS) for 0.5 hour before incubating with primary antibodies diluted in IF buffer for 1 hour. The slides were then washed 3 times with PBS and incubated with secondary antibodies (1:1000) diluted in IF buffer for 0.5 hour. The slides were subsequently washed 3 times with PBS and mounted with coverslips. All images were taken using Olympus IX71 microscope. All secondary antibodies (FITC or TRITC-labeled goat-anti-mouse/rabbit IgG) were from Jackson ImmunoResearch. Primary antibodies were used at 1:1000 dilutions unless otherwise indicated. The following antibodies were previously described: ATR⁵¹, TopBP1⁵², RAP80, BRCA1³³, CCDC98 (1:100)³², pCCDC98 (1:100)³², MERIT40 (1:100)⁵³, 53BP1¹⁸, RAD51⁵⁴, and BACH1 (1:100)³⁰. The following antibodies were purchased: ubiquitin (FK2) (Millipore, 04–263), SUMO1 (1:200, Invitrogen, 33–2400), γ H2AX (Millipore, 05–636), RAD18 (1:500, Proteintech, 18333-1-AP), pATM (1:200, Cell Signaling, 4526), rabbit against SYCP3 (Novus, NB300-232), and mouse against SYCP3 (Abcam, ab97672). The following antibodies were kind gifts from others: K48-Ub (Genentech), K63-Ub (Genentech), SUMO2/3 (1:200, Michael Matunis), MDC1 (1:100, Zhenkun Lou), CTIP (1:50, Richard Baer), and pDNAPK (1:50, David Chen). Antibodies against GST (1:1000) were generated using the full length GST protein. The specificities of the following antibodies were well established: anti-ubiquitin (FK2)⁵⁵, K48-Ub⁸, and K63-Ub⁸. The specificities of the following antibodies were confirmed since they have the same staining patterns as shown in the literature using other antibodies that recognize the same proteins: anti-SYCP3⁴⁷, SUMO1⁵⁰, SUMO2/3⁵⁰, γ H2AX³, MDC1⁴, RAD18¹⁶, ATR⁴⁷, TopBP1⁵⁶, BRCA1⁴⁷, and RAD51⁵⁷. The specificities of the following antibodies were confirmed by loss of signals in the knockouts: anti-53BP1¹⁸, RAP80³³, pATM⁵⁸, and pDNAPK⁵⁹. The specificity of anti-CTIP antibody was confirmed previously by shRNA-mediated knockdown in mouse cells⁶⁰. The specificities of the following antibodies in detecting mouse proteins in the XY body were also confirmed by using additional antibodies that recognize the same proteins: anti-pATM (Cell Signaling, 5883), pDNAPK (Abcam, ab18356), and CTIP³⁴. The specificities of anti-CCDC98 and pCCDC98 antibodies were confirmed since they recognize the same protein and have the same staining patterns in the XY body. The specificity of anti-CCDC98 antibody was further confirmed by loss of signals in the *Rap80* knockout spermatocytes. The specificities of anti-MERIT40 was confirmed by the observation that recombinant MERIT40 localized to the unsynapsed axes using the *in vitro* binding method developed in this study. The absence of BACH1 in the XY body was also confirmed by another antibody that recognizes a different region of mouse BACH1 (Santa Cruz, sc-365708).

Surface spread of spermatocytes

Spermatocytes were dissociated from seminiferous tubules by collagenase treatment (1mg/ml) at 37 °C. Isolated cells were collected and incubated for 10 minutes in buffer containing 30 nM Tris-HCl pH 8.2, 50 mM sucrose, and 17 mM sodium citrate. The cells were then pelleted and resuspended in 100 mM sucrose before being spread onto slides pre-coated with PFA solution (1% PFA, 0.15% Triton-X-100, 10 mM sodium borate, pH 9.2).

The slides were kept in a humidified chamber for 4 hours, washed with 0.4% Photoflo (Kodak), and dried.

***In vitro* binding of recombinant proteins to spermatocytes**

GST-tagged proteins were expressed in BL21 bacteria, purified using glutathione sepharose (GE), and dialyzed with 1×PBS. 10 nM of recombinant protein was added to the slides with surface spread of spermatocytes, and incubated for 1 hour in room temperature. The excess protein was then removed and the slides were washed 3 times with 1×PBS before subjected to immunofluorescence staining using antibodies against GST. To remove phosphate groups from proteins in the surface spread of spermatocytes, each slide was treated with 1000 units of lambda protein phosphatase (NEB) for 1 hour at room temperature in the buffer provided by the supplier.

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References

1. Mahadevaiah SK, et al. Recombinational DNA double-strand breaks in mice precede synapsis. *Nat Genet.* 2001; 27:271–276. [PubMed: 11242108]
2. Turner JM. Meiotic sex chromosome inactivation. *Development.* 2007; 134:1823–1831. [PubMed: 17329371]
3. Fernandez-Capetillo O, et al. H2AX is required for chromatin remodeling and inactivation of sex chromosomes in male mouse meiosis. *Dev Cell.* 2003; 4:497–508. [PubMed: 12689589]
4. Ichijima Y, et al. MDC1 directs chromosome-wide silencing of the sex chromosomes in male germ cells. *Genes Dev.* 2011; 25:959–971. [PubMed: 21536735]
5. Huen MS, et al. RNF8 transduces the DNA-damage signal via histone ubiquitylation and checkpoint protein assembly. *Cell.* 2007; 131:901–914. [PubMed: 18001825]
6. Mailand N, et al. RNF8 ubiquitylates histones at DNA double-strand breaks and promotes assembly of repair proteins. *Cell.* 2007; 131:887–900. [PubMed: 18001824]
7. Lu LY, et al. RNF8-dependent histone modifications regulate nucleosome removal during spermatogenesis. *Dev Cell.* 2010; 18:371–384. [PubMed: 20153262]
8. Newton K, et al. Ubiquitin chain editing revealed by polyubiquitin linkage-specific antibodies. *Cell.* 2008; 134:668–678. [PubMed: 18724939]
9. Wang B, Elledge SJ. Ubc13/Rnf8 ubiquitin ligases control foci formation of the Rap80/Abraxas/Brca1/Brcc36 complex in response to DNA damage. *Proc Natl Acad Sci U S A.* 2007; 104:20759–20763. [PubMed: 18077395]
10. Meerang M, et al. The ubiquitin-selective segregase VCP/p97 orchestrates the response to DNA double-strand breaks. *Nat Cell Biol.* 2011; 13:1376–1382. [PubMed: 22020440]
11. Feng L, Chen J. The E3 ligase RNF8 regulates KU80 removal and NHEJ repair. *Nat Struct Mol Biol.* 2012; 19:201–206. [PubMed: 22266820]
12. Morris JR, et al. The SUMO modification pathway is involved in the BRCA1 response to genotoxic stress. *Nature.* 2009; 462:886–890. [PubMed: 20016594]
13. Galanty Y, et al. Mammalian SUMO E3-ligases PIAS1 and PIAS4 promote responses to DNA double-strand breaks. *Nature.* 2009; 462:935–939. [PubMed: 20016603]
14. La Salle S, Sun F, Zhang XD, Matunis MJ, Handel MA. Developmental control of sumoylation pathway proteins in mouse male germ cells. *Dev Biol.* 2008; 321:227–237. [PubMed: 18602382]

15. Huang J, et al. RAD18 transmits DNA damage signalling to elicit homologous recombination repair. *Nat Cell Biol.* 2009; 11:592–603. [PubMed: 19396164]
16. van der Laan R, et al. Ubiquitin ligase Rad18Sc localizes to the XY body and to other chromosomal regions that are unpaired and transcriptionally silenced during male meiotic prophase. *J Cell Sci.* 2004; 117:5023–5033. [PubMed: 15383616]
17. Sun J, et al. Rad18 is required for long-term maintenance of spermatogenesis in mouse testes. *Mech Dev.* 2009; 126:173–183. [PubMed: 19068231]
18. Ward IM, Minn K, van Deursen J, Chen J. p53 Binding protein 53BP1 is required for DNA damage responses and tumor suppression in mice. *Mol Cell Biol.* 2003; 23:2556–2563. [PubMed: 12640136]
19. Inagaki A, et al. Meiotic functions of RAD18. *J Cell Sci.* 2011; 124:2837–2850. [PubMed: 21807948]
20. Baart EB, de Rooij DG, Keegan KS, de Boer P. Distribution of Atr protein in primary spermatocytes of a mouse chromosomal mutant: a comparison of preparation techniques. *Chromosoma.* 2000; 109:139–147. [PubMed: 10855505]
21. Wang J, Gong Z, Chen J. MDC1 collaborates with TopBP1 in DNA replication checkpoint control. *J Cell Biol.* 2011; 193:267–273. [PubMed: 21482717]
22. Wang B, et al. Abraxas and RAP80 form a BRCA1 protein complex required for the DNA damage response. *Science.* 2007; 316:1194–1198. [PubMed: 17525340]
23. Sobhian B, et al. RAP80 targets BRCA1 to specific ubiquitin structures at DNA damage sites. *Science.* 2007; 316:1198–1202. [PubMed: 17525341]
24. Kim H, Chen J, Yu X. Ubiquitin-binding protein RAP80 mediates BRCA1-dependent DNA damage response. *Science.* 2007; 316:1202–1205. [PubMed: 17525342]
25. Shao G, et al. MERIT40 controls BRCA1-Rap80 complex integrity and recruitment to DNA double-strand breaks. *Genes Dev.* 2009; 23:740–754. [PubMed: 19261746]
26. Feng L, Huang J, Chen J. MERIT40 facilitates BRCA1 localization and DNA damage repair. *Genes Dev.* 2009; 23:719–728. [PubMed: 19261748]
27. Wang B, Hurov K, Hofmann K, Elledge SJ. NBA1, a new player in the Brca1 A complex, is required for DNA damage resistance and checkpoint control. *Genes Dev.* 2009; 23:729–739. [PubMed: 19261749]
28. Huen MS, Sy SM, Chen J. BRCA1 and its toolbox for the maintenance of genome integrity. *Nat Rev Mol Cell Biol.* 2010; 11:138–148. [PubMed: 20029420]
29. Sy SM, et al. Critical roles of ring finger protein RNF8 in replication stress responses. *J Biol Chem.* 2011; 286:22355–22361. [PubMed: 21558560]
30. Yu X, Chini CC, He M, Mer G, Chen J. The BRCT domain is a phospho-protein binding domain. *Science.* 2003; 302:639–642. [PubMed: 14576433]
31. Kim H, Huang J, Chen J. CCDC98 is a BRCA1-BRCT domain-binding protein involved in the DNA damage response. *Nat Struct Mol Biol.* 2007; 14:710–715. [PubMed: 17643122]
32. Liu Z, Wu J, Yu X. CCDC98 targets BRCA1 to DNA damage sites. *Nat Struct Mol Biol.* 2007; 14:716–720. [PubMed: 17643121]
33. Wu J, Liu C, Chen J, Yu X. RAP80 protein is important for genomic stability and is required for stabilizing BRCA1-A complex at DNA damage sites in vivo. *J Biol Chem.* 2012; 287:22919–22926. [PubMed: 22539352]
34. Yu X, Chen J. DNA damage-induced cell cycle checkpoint control requires CtIP, a phosphorylation-dependent binding partner of BRCA1 C-terminal domains. *Mol Cell Biol.* 2004; 24:9478–9486. [PubMed: 15485915]
35. Yu X, Fu S, Lai M, Baer R, Chen J. BRCA1 ubiquitinates its phosphorylation-dependent binding partner CtIP. *Genes Dev.* 2006; 20:1721–1726. [PubMed: 16818604]
36. Chen PL, et al. Inactivation of CtIP leads to early embryonic lethality mediated by G1 restraint and to tumorigenesis by haploid insufficiency. *Mol Cell Biol.* 2005; 25:3535–3542. [PubMed: 15831459]
37. Lou Z, et al. MDC1 regulates DNA-PK autophosphorylation in response to DNA damage. *J Biol Chem.* 2004; 279:46359–46362. [PubMed: 15377652]

38. Wu J, et al. Chfr and RNF8 synergistically regulate ATM activation. *Nat Struct Mol Biol.* 2011; 18:761–768. [PubMed: 21706008]
39. Shanbhag NM, Rafalska-Metcalf IU, Balane-Bolivar C, Janicki SM, Greenberg RA. ATM-dependent chromatin changes silence transcription in cis to DNA double-strand breaks. *Cell.* 2010; 141:970–981. [PubMed: 20550933]
40. Celeste A, et al. Histone H2AX phosphorylation is dispensable for the initial recognition of DNA breaks. *Nat Cell Biol.* 2003; 5:675–679. [PubMed: 12792649]
41. Yuan J, Chen J. MRE11-RAD50-NBS1 complex dictates DNA repair independent of H2AX. *J Biol Chem.* 2010; 285:1097–1104. [PubMed: 19910469]
42. Fukuda T, Daniel K, Wojtasz L, Toth A, Hoog C. A novel mammalian HORMA domain-containing protein, HORMAD1, preferentially associates with unsynapsed meiotic chromosomes. *Exp Cell Res.* 2010; 316:158–171. [PubMed: 19686734]
43. Wojtasz L, et al. Mouse HORMAD1 and HORMAD2, two conserved meiotic chromosomal proteins, are depleted from synapsed chromosome axes with the help of TRIP13 AAA-ATPase. *PLoS Genet.* 2009; 5:e1000702. [PubMed: 19851446]
44. Shin YH, et al. Hormad1 mutation disrupts synaptonemal complex formation, recombination, and chromosome segregation in mammalian meiosis. *PLoS Genet.* 2010; 6:e1001190. [PubMed: 21079677]
45. Kogo H, et al. HORMAD1-dependent checkpoint/surveillance mechanism eliminates asynaptic oocytes. *Genes Cells.* 2012; 17:439–454. [PubMed: 22530760]
46. Wojtasz L, et al. Meiotic DNA double-strand breaks and chromosome asynapsis in mice are monitored by distinct HORMAD2-independent and -dependent mechanisms. *Genes Dev.* 2012; 26:958–973. [PubMed: 22549958]
47. Turner JM, et al. BRCA1, histone H2AX phosphorylation, and male meiotic sex chromosome inactivation. *Curr Biol.* 2004; 14:2135–2142. [PubMed: 15589157]
48. Turner JM, et al. Silencing of unsynapsed meiotic chromosomes in the mouse. *Nat Genet.* 2005; 37:41–47. [PubMed: 15580272]
49. Rogakou EP, Boon C, Redon C, Bonner WM. Megabase chromatin domains involved in DNA double-strand breaks in vivo. *J Cell Biol.* 1999; 146:905–916. [PubMed: 10477747]
50. Vigodner M. Sumoylation precedes accumulation of phosphorylated H2AX on sex chromosomes during their meiotic inactivation. *Chromosome Res.* 2009; 17:37–45. [PubMed: 19156530]
51. Ward IM, Minn K, Chen J. UV-induced ataxia-telangiectasia-mutated and Rad3-related (ATR) activation requires replication stress. *J Biol Chem.* 2004; 279:9677–9680. [PubMed: 14742437]
52. Yamane K, Wu X, Chen J. A DNA damage-regulated BRCT-containing protein, TopBP1, is required for cell survival. *Mol Cell Biol.* 2002; 22:555–566. [PubMed: 11756551]
53. Bian C, Wu R, Cho K, Yu X. Loss of BRCA1-A Complex Function in RAP80 Null Tumor Cells. *PLoS One.* 2012; 7:e40406. [PubMed: 22792303]
54. Zhang F, Wu J, Yu X. Integrator3, a partner of single-stranded DNA-binding protein 1, participates in the DNA damage response. *J Biol Chem.* 2009; 284:30408–30415. [PubMed: 19759019]
55. Fujimuro M, Sawada H, Yokosawa H. Production and characterization of monoclonal antibodies specific to multi-ubiquitin chains of polyubiquitinated proteins. *FEBS Lett.* 1994; 349:173–180. [PubMed: 7519568]
56. Barchi M, et al. Surveillance of different recombination defects in mouse spermatocytes yields distinct responses despite elimination at an identical developmental stage. *Mol Cell Biol.* 2005; 25:7203–7215. [PubMed: 16055729]
57. Moens PB, et al. Rad51 immunocytology in rat and mouse spermatocytes and oocytes. *Chromosoma.* 1997; 106:207–215. [PubMed: 9254722]
58. Dar I, Biton S, Shiloh Y, Barzilai A. Analysis of the ataxia telangiectasia mutated-mediated DNA damage response in murine cerebellar neurons. *J Neurosci.* 2006; 26:7767–7774. [PubMed: 16855104]
59. Chan DW, et al. Autophosphorylation of the DNA-dependent protein kinase catalytic subunit is required for rejoining of DNA double-strand breaks. *Genes Dev.* 2002; 16:2333–2338. [PubMed: 12231622]

60. Zhang Y, Jasin M. An essential role for CtIP in chromosomal translocation formation through an alternative end-joining pathway. *Nat Struct Mol Biol.* 2011; 18:80–84. [PubMed: 21131978]

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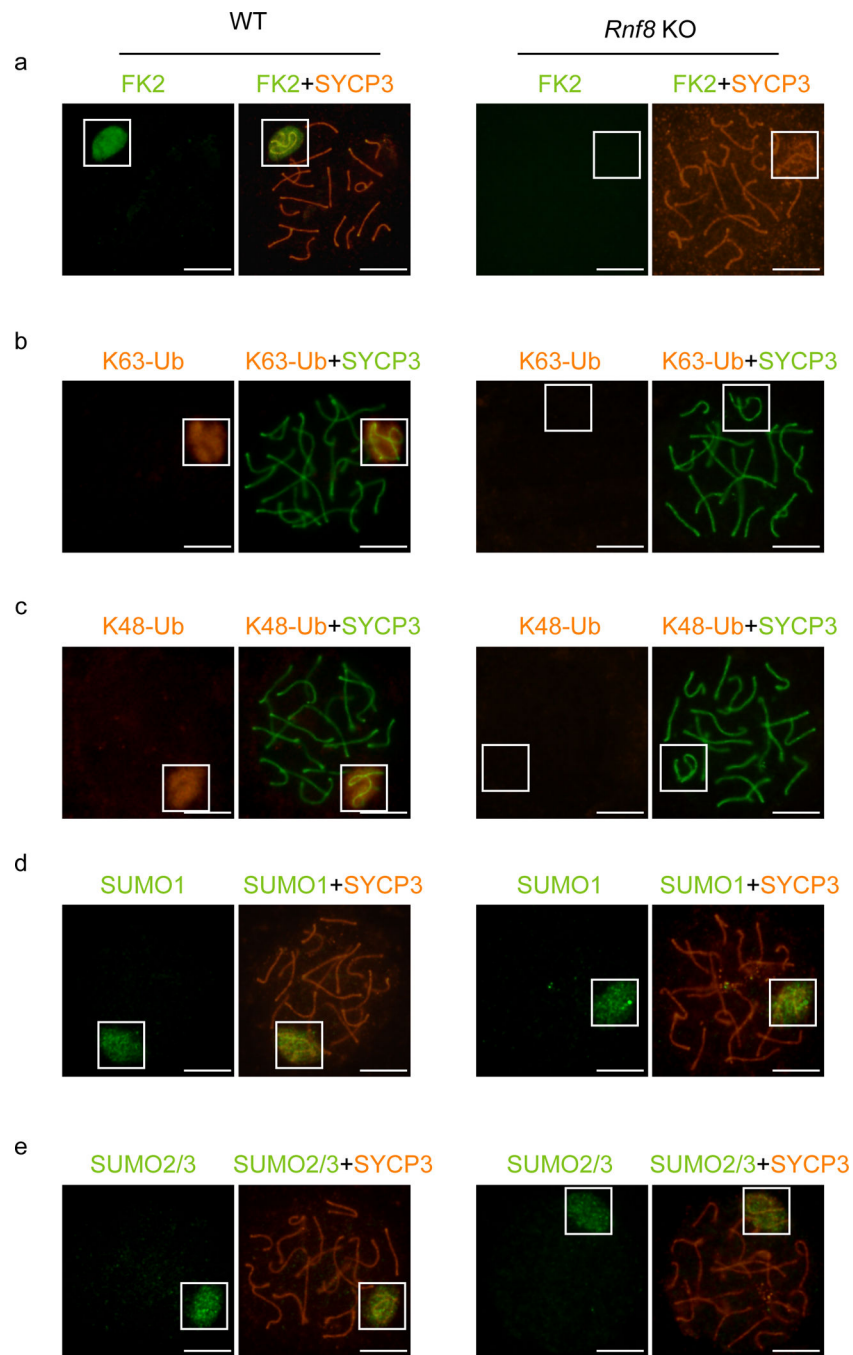


Figure 1. RNF8 controls protein ubiquitination but not sumoylation in the XY body
a.-e. Spermatocytes at pachynema from WT and *Rnf8* knockout mice were stained using antibodies against ubiquitin-FK2 (a), K63-linked ubiquitin chains (b), K48-linked ubiquitin chains (c), SUMO1 (d), and SUMO2/3 (e). SYCP3 was used to visualize the chromosome synapsis and to differentiate spermatocytes at various stages of meiotic prophase. Scale bar, 10 μ M.

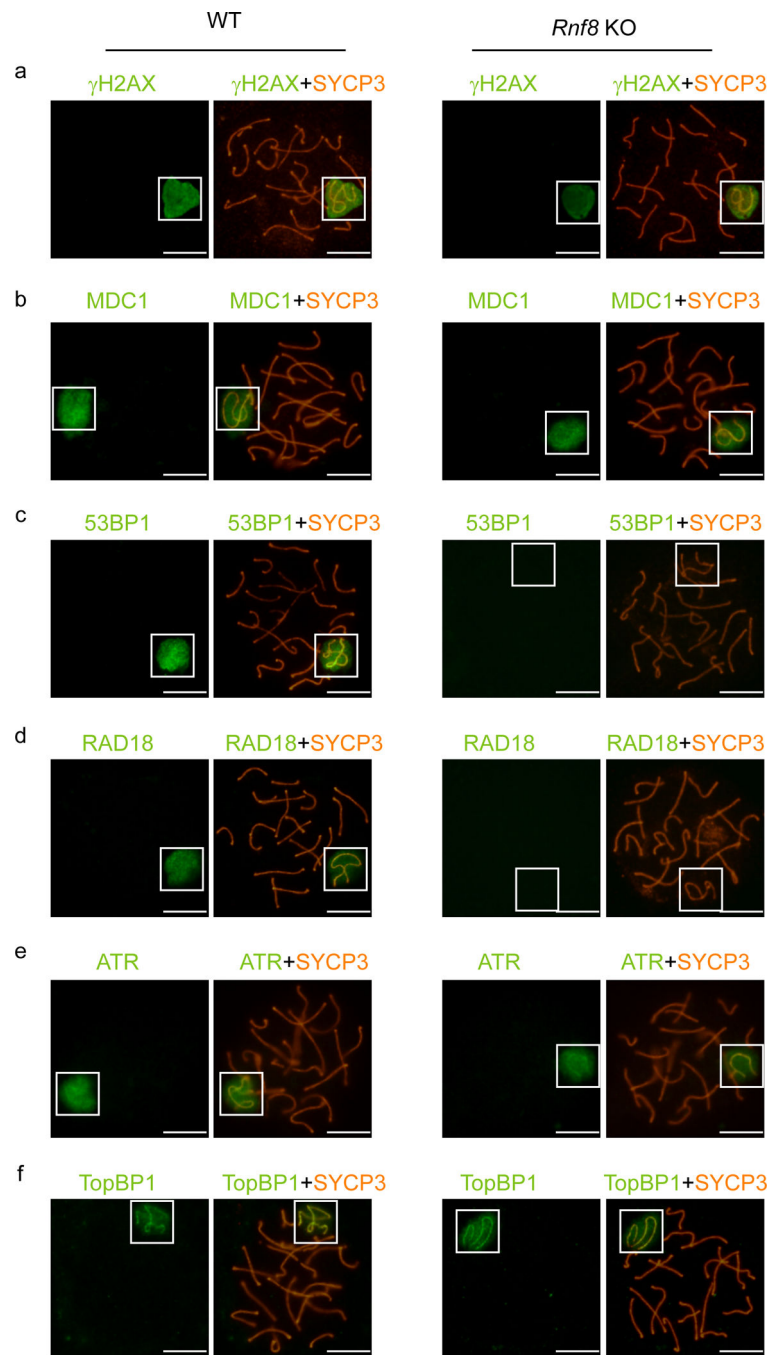


Figure 2. RNF8 controls the localization of DDR proteins over the sex chromosomes
a.-f. Spermatocytes at pachynema from WT and *Rnf8* knockout mice were stained using the antibodies antibodies against γ H2AX (a), MDC1 (b), 53BP1 (c), RAD18 (d), ATR (e), and TopBP1 (f). SYCP3 was used to visualize the chromosome synapsis and to differentiate spermatocytes at various stages of meiotic prophase. Scale bar, 10 μ M.

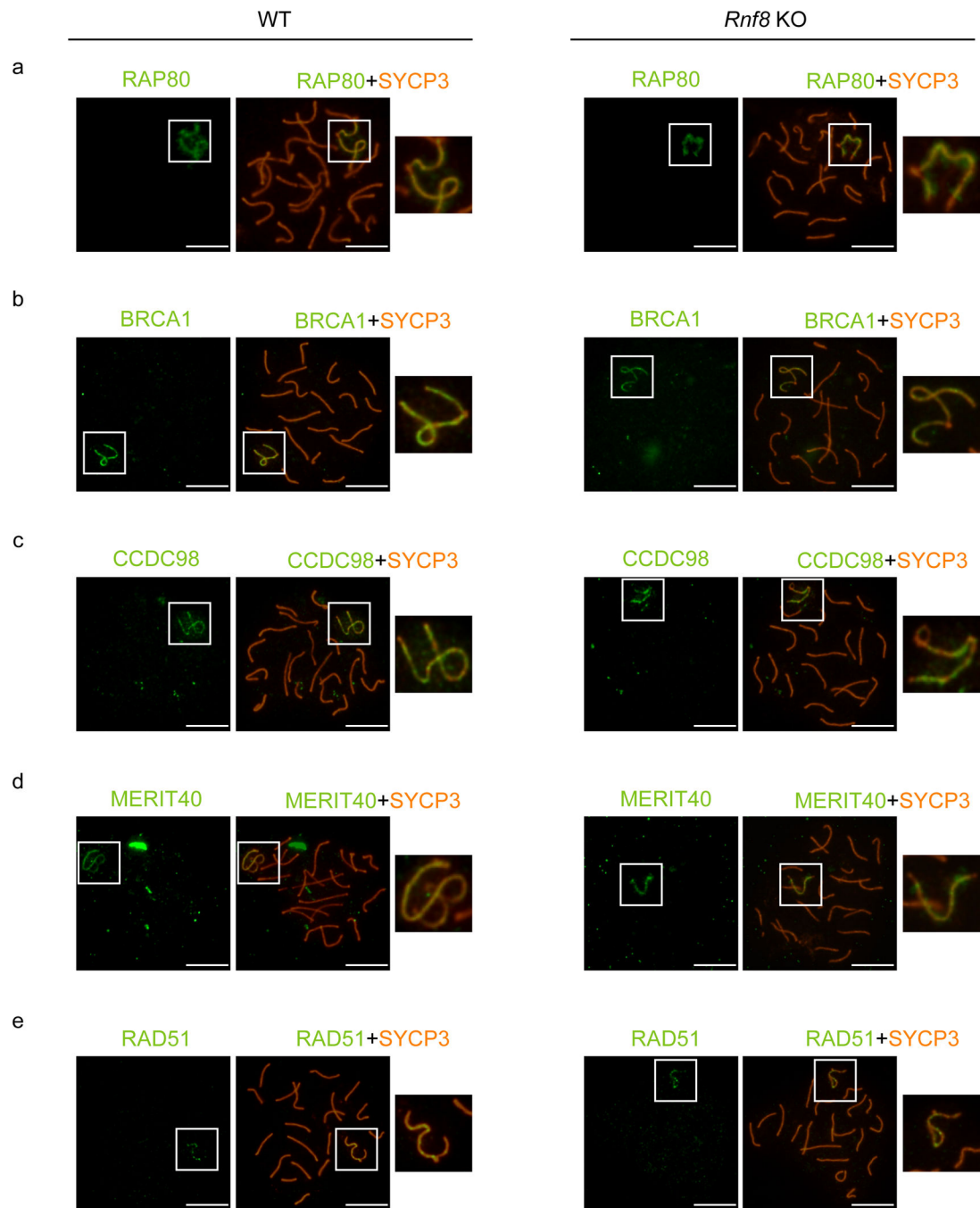


Figure 3. RNF8 is dispensable for the localization of the BRCA1-A complex

a.-e. Spermatocytes at pachynema from WT and *Rnf8* knockout mice were stained using the antibodies against RAP80 (a), BRCA1 (b), CCDC98 (c), MERIT40 (d), and RAD51 (e). SYCP3 was used to visualize the chromosome synapsis and to differentiate spermatocytes at various stages of meiotic prophase. White boxes indicate the positions of the unsynapsed axes of sex chromosomes, and are magnified on the right. Scale bar, 10 μ M.

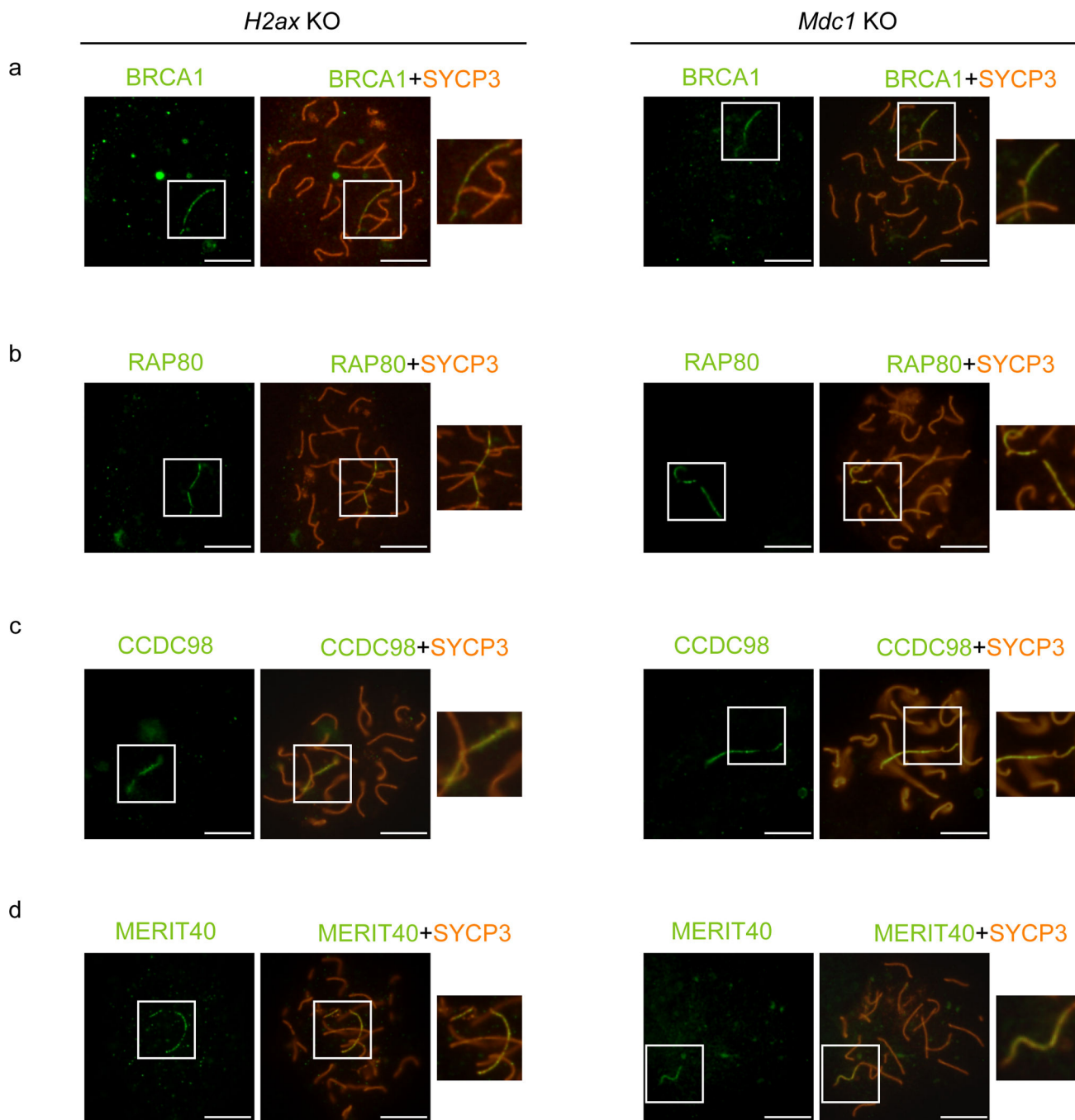


Figure 4. H2AX and MDC1 are dispensable for the localization of the BRCA1-A complex
 a.-d. Spermatocytes at pachynema from *H2ax* and *Mdc1* knockout mice were stained using the antibodies against BRCA1 (a), RAP80 (b), CCDC98 (c), and MERIT40 (d). SYCP3 was used to visualize the chromosome synapsis and to differentiate spermatocytes at various stages of meiotic prophase. White boxes indicate the positions of the unsynapsed axes of sex chromosomes, and are magnified on the right. Scale bar, 10 μ M.

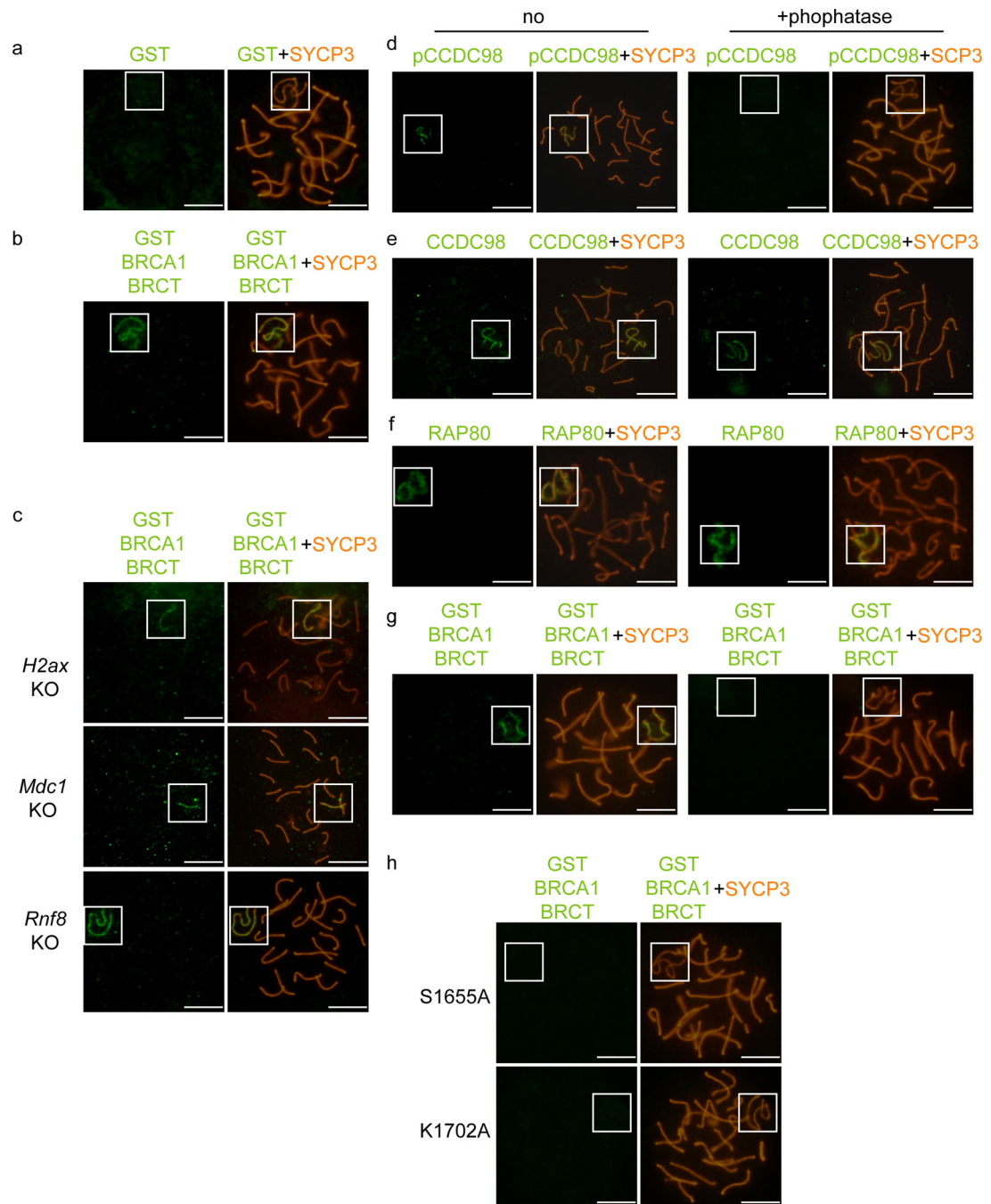


Figure 5. The localization of BRCA1 relies on BRCT domain recognition

a.-b. Surface spreads of spermatocytes from WT mice were incubated with GST-tagged recombinant proteins as indicated, and stained using antibodies against GST. c. Surface spreads of spermatocytes from *H2ax*, *Mdc1*, and *Rnf8* knockout mice were incubated with GST-tagged BRCA1 BRCT domain proteins, and stained using antibodies against GST. d.-f. Surface spreads of spermatocytes from WT mice were treated with or without lambda protein phosphatase (phosphatase), and stained using the antibodies as indicated. g. Surface spreads of spermatocytes from WT mice were treated with or without lambda protein

phosphatase (phosphatase), incubated with GST-tagged BRCA1 BRCT domain proteins, and stained using antibodies against GST. h. Surface spreads of spermatocytes from WT mice were incubated with GST-tagged recombinant proteins as indicated, and stained using antibodies against GST. In all panels, SYCP3 was used to visualize the chromosome synapsis and to differentiate spermatocytes at various stages of meiotic prophase. White boxes indicate the positions of the unsynapsed axes of sex chromosomes. Scale bar, 10 μ M.

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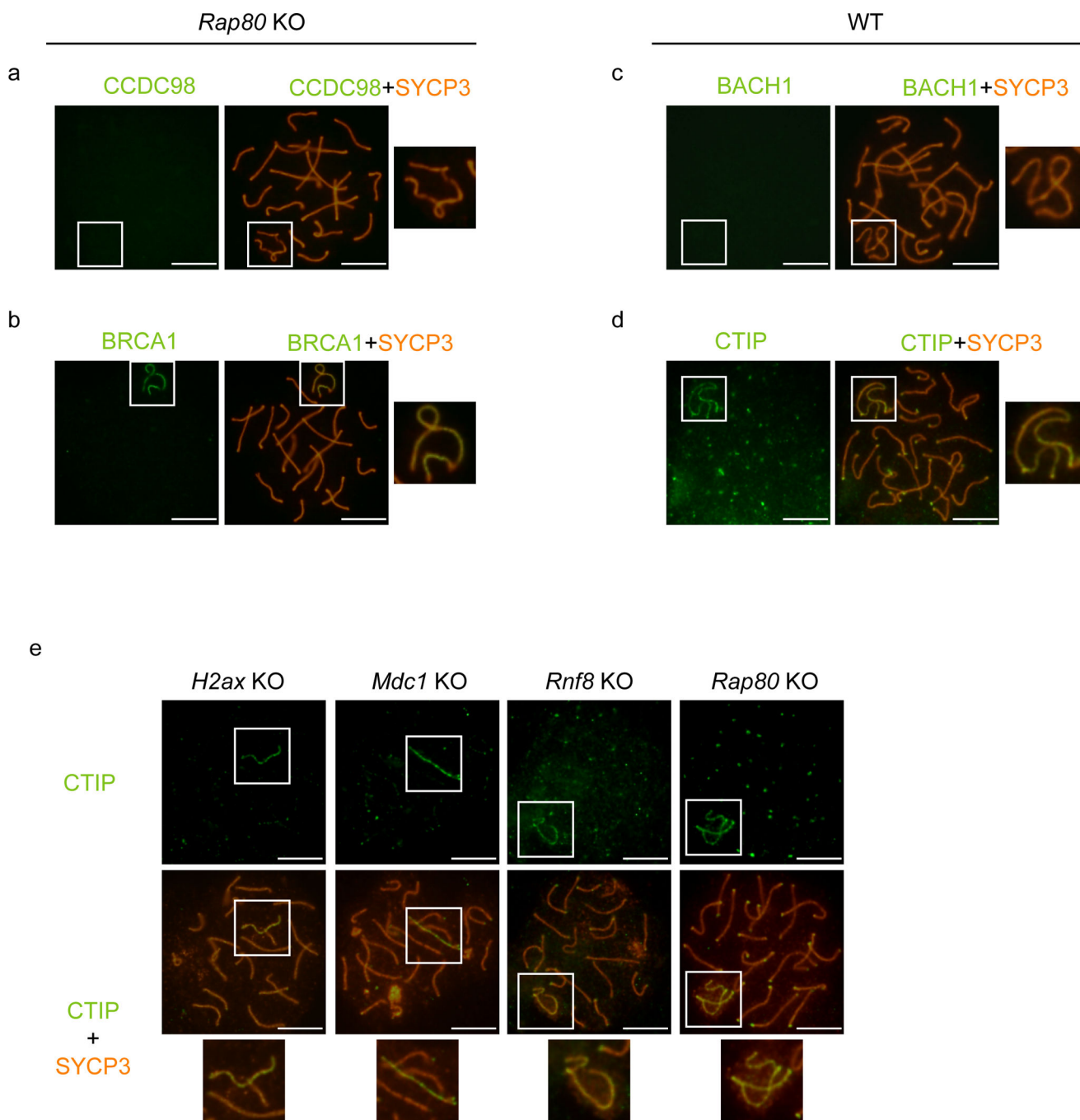


Figure 6. The localization of BRCA1 does not require the BRCA1-A complex

a.-e. Spermatocytes at pachynema from WT and *H2ax*, *Mdc1*, *Rnf8*, and *Rap80* knockout mice were stained using the antibodies against CCDC98 (a), BRCA1 (b), BACH1 (c), and CTIP (d-e). SYCP3 was used to visualize the chromosome synapsis and to differentiate spermatocytes at various stages of meiotic prophase. White boxes indicate the positions of the unsynapsed axes of sex chromosomes, and are magnified on the right. Scale bar, 10 μM.

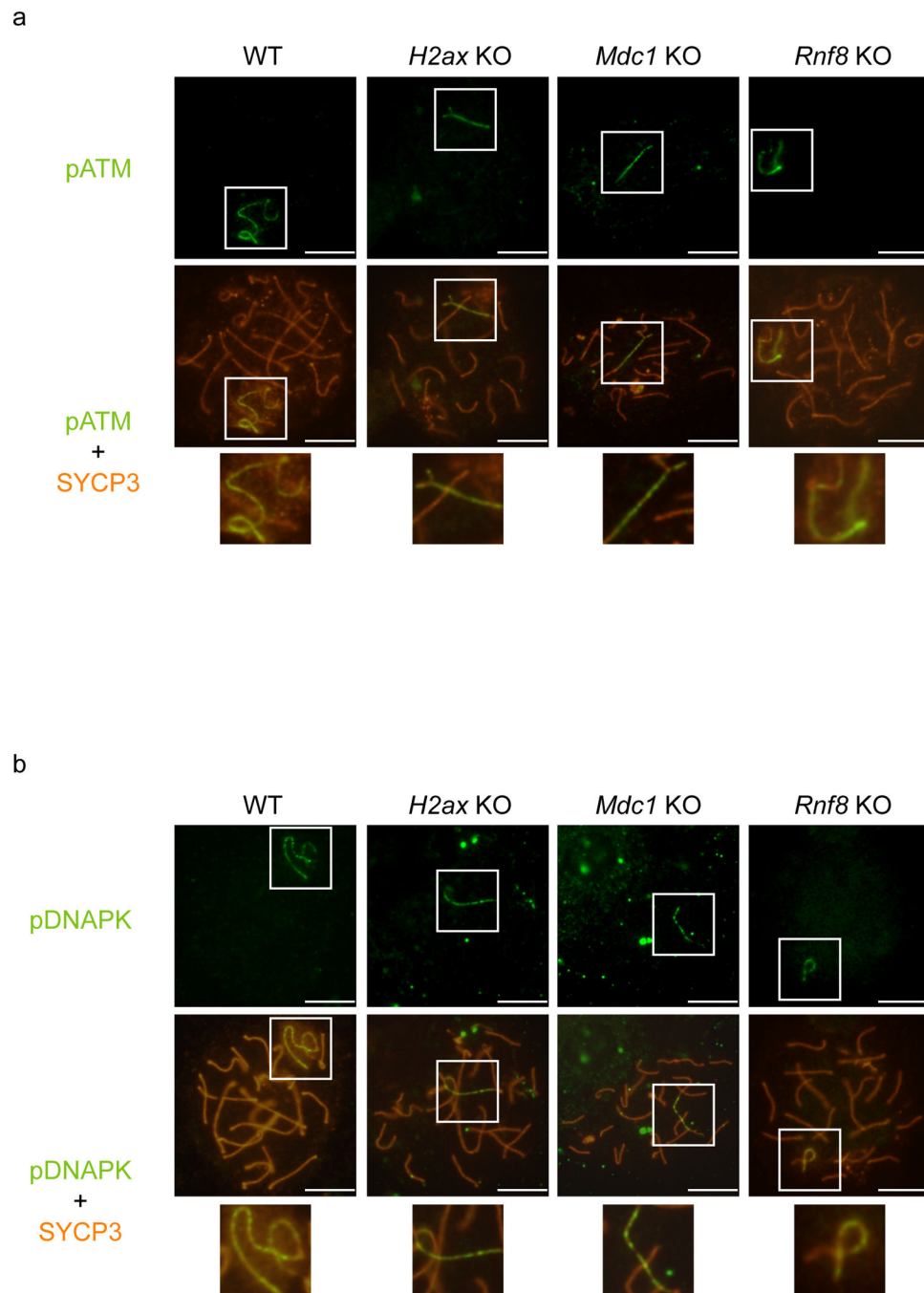
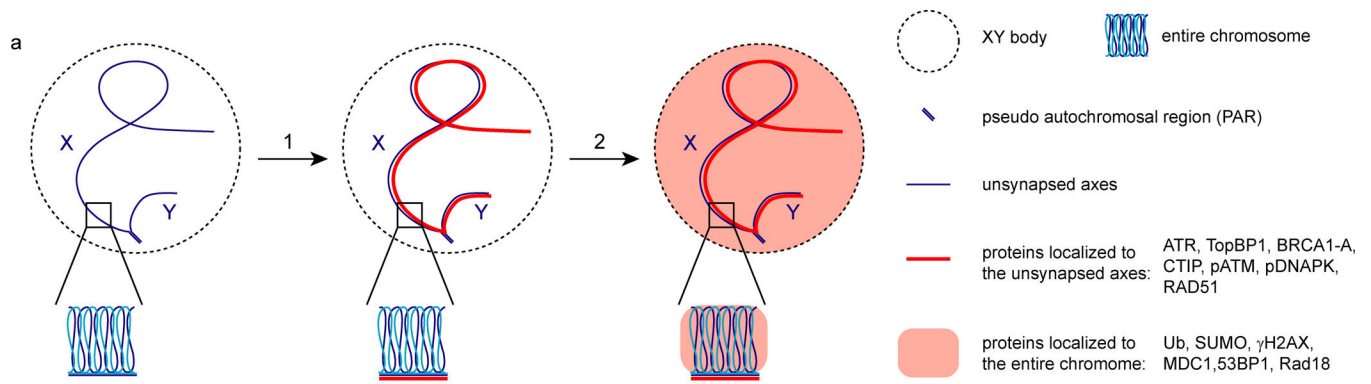


Figure 7. ATM and DNAPK localize to the unsynapsed axes of sex chromosomes

a.-b. Spermatocytes at pachynema from WT and *H2ax*, *Mdc1*, and *Rnf8* knockout mice were stained using the antibodies against activated phosphorylated ATM (a) and activated phosphorylated DNAPK (b). SYCP3 was used to visualize the chromosome synapsis and to differentiate spermatocytes at various stages of meiotic prophase. White boxes indicate the positions of the unsynapsed axes of sex chromosomes, and are magnified on the bottom. Scale bar, 10 μ M.



b

DDR proteins	XY body								Somatic IRIF			
	WT		ref	<i>Rnf8</i> KO		<i>H2ax1Mdc1</i> KO		ref	WT	<i>Rnf8</i> KO	<i>H2ax1Mdc1</i> KO	ref
	axes	chomo-somes		axes	chomo-somes	axes	chomo-somes					
K48/63-Ubiquitination	✓	✓		X	X	X	X	3,4	✓	X	X	9-11
Sumoylation	✓	✓	14	✓	✓	✓	✓	3,4	✓	X	X	12,13
53BP1	✓	✓		X	X	X	X	3,4	✓	X, early	X, early	41
RAD18	✓	✓	16	X	X	X	X	3,4	✓	X	X	15
ATR	✓	✓	20	✓	✓	✓	X	3,4	?	?	?	
TopBP1	✓	✓	56	✓	✓	✓	X	3,4	✓	✓	X	21
RAD51	✓	X	57	✓	X	✓	X	3,4	✓	↓	↓	29
BRCA1-A	✓	X		✓	X	✓	X		✓	X, early	X, early	40
CTIP	✓	X		✓	X	✓	X		✓	?	?	35

✓: presence X: absence ↓: decreased ?: unknown

H2AX and MDC1 KO are combined due to the shared properties among these two cells.

Early refers to IRIF within 15 minutes.

The shaded boxes are localization patterns found in this study. The others are derived from literature data. (ref: reference)

Figure 8. DDR at the XY body and at somatic DNA damage sites are different

a. Model of the DNA damage response in the XY body. Some proteins are directly recruited to the unsynapsed axes independent of canonical DDR pathways. The DDR then spreads over the sex chromosomes by amplification of γ H2AX-MDC1-RNF8 signaling, which recruits another set of proteins. b. Summary of the localizations of DDR proteins in XY body at meiotic prophase and their formations of ionizing radiation-induced foci (IRIF) in somatic cells.